<u>AMENDMENTS</u>

In the Claims

Claims 32-43 are pending.

Claim 41 is amended as follows:

Claims 1-31 (canceled).

- 32. (previously presented) A method of diagnosing colorectal cancer comprising:
- a) determining the expression of a nucleic acid sequence at least 95% identical to SEQ ID No:1 in a first sample comprising colorectal cancer cells from a first individual; and
- b) comparing the expression of said nucleic acid sequence in the first sample to expression of said nucleic acid sequence in a second sample taken from either normal colon tissue of said first individual or from normal colon tissue, or a bodily secretion of a second unaffected individual;

wherein a difference between the expression in the first sample and the expression in the second sample indicates that the first individual has colorectal cancer.

- 33. (previously presented) The method of claim 32, wherein said second sample is from said first individual.
- 34. (previously presented) The method of claim 33, wherein said first sample is colorectal tissue.
- 35. (previously presented) The method of claim 33, wherein said second sample is colorectal tissue.
- 36. (previously presented) The method of claim 33, wherein said second sample is cancerous tissue.

- 37. (previously presented) The method of claim 32, wherein said second sample is from a second individual.
- 38. (previously presented) The method of claim 37, wherein said first sample is colorectal tissue.
- 39. (previously presented) The method of claim 37, wherein said second sample is colorectal tissue.
- 40. (previously presented) The method of claim 37, wherein said second sample is cancerous tissue.
- 41. (currently amended) The method of claim 32, wherein said gene nucleic acid sequence is SEQ ID NO:1.
- 42. (previously presented) The method of claim 32, wherein said expression is measured using a labeled nucleic acid probe.
- 43. (previously presented) The method of claim 32, wherein said expression is measured utilizing a biochip.

Claims 44-47 (canceled).